

Rhesus monkey brain development during late infancy and the effect of phencyclidine: A longitudinal MRI and DTI study



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ARTICLE INFO

Article history:

Accepted 27 November 2014

Available online 5 December 2014

Keywords:

Infant
Neurodevelopment
Primates
Template
Phencyclidine

ABSTRACT

Early brain development is a complex and rapid process, the disturbance of which may cause the onset of brain disorders. Based on longitudinal imaging data acquired from 6 to 16 months postnatal, we describe a systematic trajectory of monkey brain development during late infancy, and demonstrate the influence of phencyclidine (PCP) on this trajectory. Although the general developmental trajectory of the monkey brain was close to that of the human brain, the development in monkeys was faster and regionally specific. Gray matter volume began to decrease during late infancy in monkeys, much earlier than in humans in whom it occurs in adolescence. Additionally, the decrease of gray matter volume in higher-order association regions (the frontal, parietal and temporal lobes) occurred later than in regions for primary functions (the occipital lobe and cerebellum). White matter volume displayed an increasing trend in most brain regions, but not in the occipital lobe, which had a stable volume. In addition, based on diffusion tensor imaging, we found an increase in fractional anisotropy and a decrease in diffusivity, which may be associated with myelination and axonal changes in white matter tracts. Meanwhile, we tested the influence of 14-day PCP treatment on the developmental trajectories. Such treatment tended to accelerate brain maturation during late infancy, although not statistically significant. These findings provide comparative information for the understanding of primate brain maturation and neurodevelopmental disorders.

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Introduction

Understanding brain development during infancy, a critical stage with rapid brain changes and vulnerabilities to exogenous influences, is essential for the research of pathological mechanisms of neurodevelopmental disorders. Recent advances in brain imaging techniques enable investigating brain maturation *in-vivo*. However, partly due to the long life span of humans, most human developmental studies are based on cross-sectional or mixed-longitudinal data that provide less accurate observations than longitudinal studies, especially when there is a large individual variability in brain development (Casey et al., 2005). Moreover, it is impractical to interfere with normal brain

development of humans, which impedes the understanding of neurodevelopmental disorders.

The rhesus monkey, a widely-used experimental animal with phylogenetic closeness to humans (Bizon and Woods, 2009), has a shorter life span than humans. Thus it is a good candidate for longitudinal studies of brain development and neurodevelopmental disorders. However, there are few comparative imaging studies on monkey brain development, and most of the existing studies are cross-sectional (Chen et al., 2013; Shi et al., 2013). A pioneering longitudinal imaging study on monkey brain development was performed 8 years ago (Malkova et al., 2006). This study investigated the growth of total brain volume and total white matter volume from birth to late puberty. Due to the limitation of the data (based on 1.5 T MRI scanners), the study did not provide important information of gray matter and regional brain growth that are essential for the characterization of the trajectories of brain development.

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On the other hand, when using monkeys as experimental animals, we can use various ways to disturb brain development. Phencyclidine (PCP), a noncompetitive N-methyl-D-aspartate (NMDA) receptor antagonist, is widely used to build animal models of schizophrenia (Jones et al., 2011; Mouri et al., 2007). Recently, schizophrenia has been regarded as a neurodevelopmental disorder, whose pathology may be contributed by disturbances during prenatal and postnatal brain maturation (Jaaro-Peled et al., 2009; Lewis and Levitt, 2002; Rapoport et al., 2005). Based on the hypothesis, previous studies have examined the effects of PCP on brain development in rodents, and they suggest that early postnatal PCP treatment is a novel promising animal model for schizophrenia (Broberg et al., 2008, 2013; Kjaerby et al., 2014; Rasmussen et al., 2007; Wang et al., 2001). However, species differences between rodents and humans constitute a major gap for “translational psychiatry”, which monkeys could help to fill.

Our study focuses on a comprehensive longitudinal analysis of monkey brain development during late infancy, and on the influence of PCP on the development by using non-invasive brain imaging techniques and linear mixed-effect models. To our knowledge, this is the first longitudinal MRI-based study that characterizes regional growth of gray and white matter, and the first longitudinal DTI-based study that characterizes white matter maturation during early brain development of monkeys, as well as the first work to investigate the influence of PCP on these developmental trajectories.

Materials and methods

Monkeys

A total of 14 male rhesus macaques (*Macaca mulatta*), from the breeding colonies at the Primate Center of the Kunming Institute of Zoology Chinese Academy of Sciences, were scanned at 5 different time-points: a baseline scan (at the age of 196 ± 8 days postnatal) and about 40, 68, 105, and 303 days later (Fig. 1A, details shown in Table S1). These scans covered the period of late infancy, which spans from ~0.44 years postnatal (the eruption of the last deciduous tooth) to ~1.35 years postnatal (the eruption of the first permanent tooth) (Smith et al., 1994). For convenience, these scans were named 6 months (S6), 7 months (S7), 8 months (S8), 10 months (S10), and 16 months (S16), respectively. In S6, S10 and S16, all 14 monkeys were scanned; and in S7 and S8, only 10 monkeys were scanned. After the first scan, 10 monkeys had a 14-day phencyclidine (PCP, Chemsky (ShangHai) International Co., Ltd. #20120516) treatment (twice per day). Of these 10 monkeys, 5 received PCP with a dose 0.15 mg/kg body weight (PCP1), and the other 5 monkeys with a dose of 0.3 mg/kg body weight (PCP2). The remaining 4 monkeys were controls treated with saline solution (0.1 ml/kg, Control). After birth, the monkeys were reared with their mothers in the breeding center. At the age of 6 months, they were moved into other cages for experiments (two monkeys per

cage), and were housed under standard conditions (a 12-hr light/dark cycle with light on from 07:00 to 19:00; humidity at 60%, temperature at 21 ± 2 °C, 5 times/day deliveries of food and milk). This is the standard breeding protocol used in our primate center to reduce the adverse influence of maternal separation and the risk of naturally occurring maternal neglect and abuse. The experiments were conducted in accordance with the guidelines for the National Care and Use of Animals approved by the National Animal Research Authority of China. Detailed scan ages and other information for each monkey are presented in Table S1.

Data acquisition

MRI and DTI data were acquired with a PHILIPS Achieva 3.0 T MR unit of The First Affiliated Hospital of Kunming Medical University, Kunming, Yunnan, China. Before the scan, monkeys were intramuscularly injected with atropine (0.05 mg/kg, ShangHai Harvest Pharmaceutical Co., LTD.). Then the monkeys were deeply anesthetized with ketamine (7.5 mg/kg, im, ShengYang Animal Pharmaceutical factory) followed by pentobarbital sodium (5 mg/kg for 1.3–1.9 kg body weight and 15 mg/kg for 2.2–3.5 kg body weight, XiTang Biological Company). The anesthesia was stable for at least 1 h, and the scan duration was less than 1 h. Thus, no complementary anesthesia was needed during the scan. A three-dimensional sagittal T1-weighted (T1-w) anatomical image was acquired using an inversion-recovery prepared 3-D spoiled gradient echo (SPGR) pulse sequence (TR = 14 ms; TE = 7 ms; flip angle = 8°; field of view = 160 mm; matrix = 320×320; number of averages = 2; final voxel resolution = $0.6 \times 0.5 \times 0.5$ mm³; scan duration = 8:30 min). DTI data were acquired using a two-dimensional echo-planar diffusion-weighted spin-echo sequence (TR = 8 s; TE = 78 ms; flip angle = 90°; field of view = 120 mm; matrix = 96 × 96; slice thickness = 1.25 mm, no gap; number of averages = 3; final voxel resolution = $1.25 \times 1.25 \times 1.25$ mm³; b = 1000 s/mm²; 32 directions with one non-diffusion weighted image; scan duration = 15 min).

Template construction

MRI (T1-w) template construction

Brain tissue was manually delineated from non-brain tissue by trained researchers for each dataset. Field inhomogeneity was bias-corrected using FAST in FSL (Smith et al., 2004; Zhang et al., 2001), which also provided segmentation maps including tissue class probability images and partial volume images. The MRI template was constructed using the Advanced Normalization Tools (ANTS) software (Avants et al., 2008; Klein et al., 2009). First, all images were rotated to the same orientation by a rigid alignment, demeaned by dividing their mean values, and then averaged to create an initial template with 0.5 mm³ isotropic resolution. The template was iteratively optimized by transforming all input images to the current template and

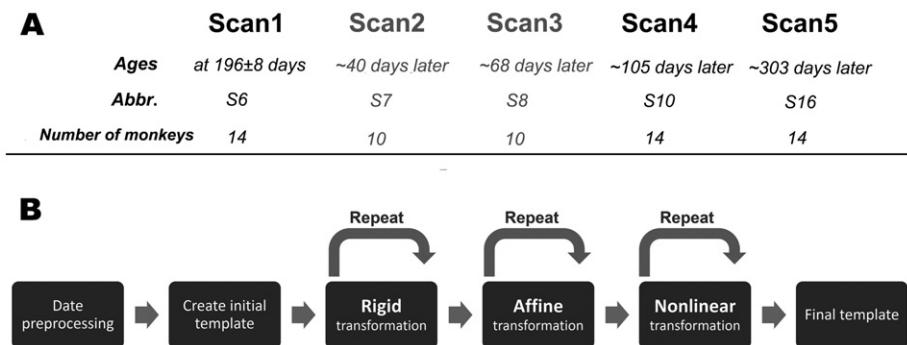


Fig. 1. Age and number of monkeys for each scan (A); and procedure of template creation (B). Starting at the second scan, dates are represented as the number of days after the first scan. The scans are abbreviated as S6 (6 months), S7 (7 months), S8 (8 months), S10 (10 months), and S16 (16 months). S6, S10 and S16 include all 14 monkeys and are used for linear mixed-model analysis. Detailed scan and age information of each monkey are presented in Table S1.

averaged to create a refined template. The procedure was performed using rigid, affine, and nonlinear alignments, and repeated until the average image converged for each type of alignment (Fig. 1B). The transforms from different alignments were concatenated so that the input images were only re-sampled once to minimize interpolation errors. After the final template was constructed, probabilistic tissue segment maps of each image were warped into the template space, smoothed with a 1-mm FWHM Gaussian smoothing kernel, and averaged to create tissue probability maps. Furthermore, brain parcellation maps were created using the following steps: 1) The template was transformed to the T1-w image of the UNC Primate Brain Atlas (Styner et al., 2007) by an affine alignment and then a nonlinear alignment. 2) The template was labeled by inverse-transforming UNC parcellation maps to the template space based on the transformations estimated from the previous alignments. 3) The template label was carefully checked and manually adjusted. The parcellation maps were used for ROI-based tissue volume analysis.

DTI template construction

Artifacts from eddy currents and motion in the diffusion data were corrected using FDT of FSL (Smith et al., 2004). For each dataset, the brain was manually delineated from non-brain tissue on the non-diffusion weighted image by trained researchers, and then the diffusion tensor was reconstructed using FSL. The DTI template was constructed using DTI-TK, a spatial normalization toolkit optimized for examining white matter morphometry based on DTI data (Zhang et al., 2007). DTI-TK uses higher order information of diffusion tensor images for

spatial normalization and has been used to generate high-quality DTI templates of adult humans (Zhang et al., 2011), human infants (Wang et al., 2011), and adult primates (Adluru et al., 2012). The procedures and parameters used to generate a monkey DTI template were similar to those employed in a previous study (Adluru et al., 2012). All input tensor images were rotated to the same orientation by a rigid alignment, and then averaged using a Log-Euclidean mean to create an initial template with a 0.5-mm³ isotropic resolution. Log-Euclidean tensor averaging preserved white matter orientation with minimal blurring. The template was iteratively optimized using rigid and then affine alignments, and repeated until the average image converged for each type of alignment. Finally, the template was iteratively optimized using nonlinear alignments.

Brain development analysis

Total brain volume

Total brain volume was estimated for each subject based on the manually de-skinned structural images, and analyzed using linear mixed-effect models (see below).

ROI-based tissue volume analysis

To avoid bias to particular time points, all structural images were transformed to the T1-w template by an affine alignment and then by a non-linear alignment. Gray matter and white matter partial volume images for each dataset were warped into the template space using the transformations from the previous alignments, and modulated by

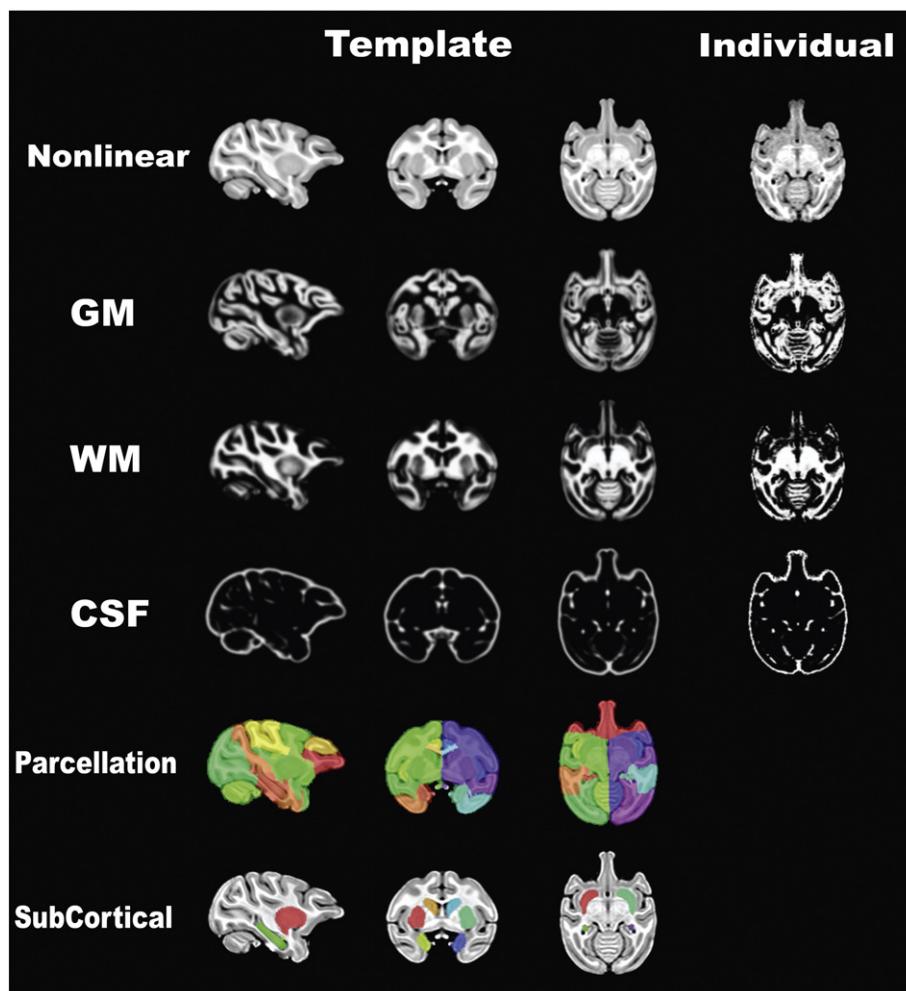


Fig. 2. MRI-based template. The first three columns illustrate the template with corresponding tissue probability maps and parcellation maps. The fourth column shows the representative images of one individual monkey.

multiplying the Jacobian determinants of the deformation field to compensate for the expansion/reduction from the nonlinear transformation (Good et al., 2001). Finally, tissue volumes for gray and white matter in each brain region were estimated based on wrapped partial volume images. Besides, the ROI-based analysis was also performed in the native space by inverse-transforming the labels of brain regions into the images of each monkey to avoid the interpolation and smoothing. The results in the native space were similar to the results in the template space.

Voxel-based morphometry (VBM)

Similar to the ROI-based analysis, gray matter and white matter partial volume images were warped into the template space and modulated by multiplying the Jacobian determinants of the deformation field. Wrapped partial volume images were smoothed with a 1-mm FWHM Gaussian smoothing kernel and used for the linear mixed-effect analysis. The analysis was restricted to a gray matter/white matter mask from the MRI template with corresponding tissue probabilities greater than 20% to exclude voxels with low tissue probabilities (mainly caused by spatial smoothing). Finally, the pre-processed images were used for linear mixed-effect analysis. Because the ROI-based volume analysis provided information similar to that by VBM, the results of VBM were complementary and presented in Fig. S1.

Deformation-based morphometry (DBM)

After the affine alignment accounting for the global brain size and shape difference, subject brain images of each time point were nonlinearly transformed to corresponding images of other time points using ANTs (Avants et al., 2008). The Jacobian determinants, which represented the local shape difference at each voxel between two time points, were then computed for the statistical analysis.

Tract-based spatial statistics (TBSS)

TBSS was an observer-independent method for group-wise comparisons of DTI-derived measures (Smith et al., 2006). TBSS comprised three steps: 1) normalizing input images to the template space; 2) creating a mean fractional anisotropy (FA) skeleton and projecting data to the skeleton; 3) statistical analysis. The first step was performed using DTI-TK (tensor-based spatial normalization) to register DTI images to the template space, because DTI-TK provided better normalization for DTI images than the FA-based method implemented in FSL (Bach et al., 2014; Zhang et al., 2007). The second step was performed using the standard TBSS protocol in FSL (Smith et al., 2004, 2006). FA images were calculated from spatial-normalized tensor images and averaged to create a mean FA image as well as a mean FA skeleton under an FA threshold over 0.2. A lower threshold (FA = 0.15) was also tested due to the low FA of infant monkeys, and the results were similar to those

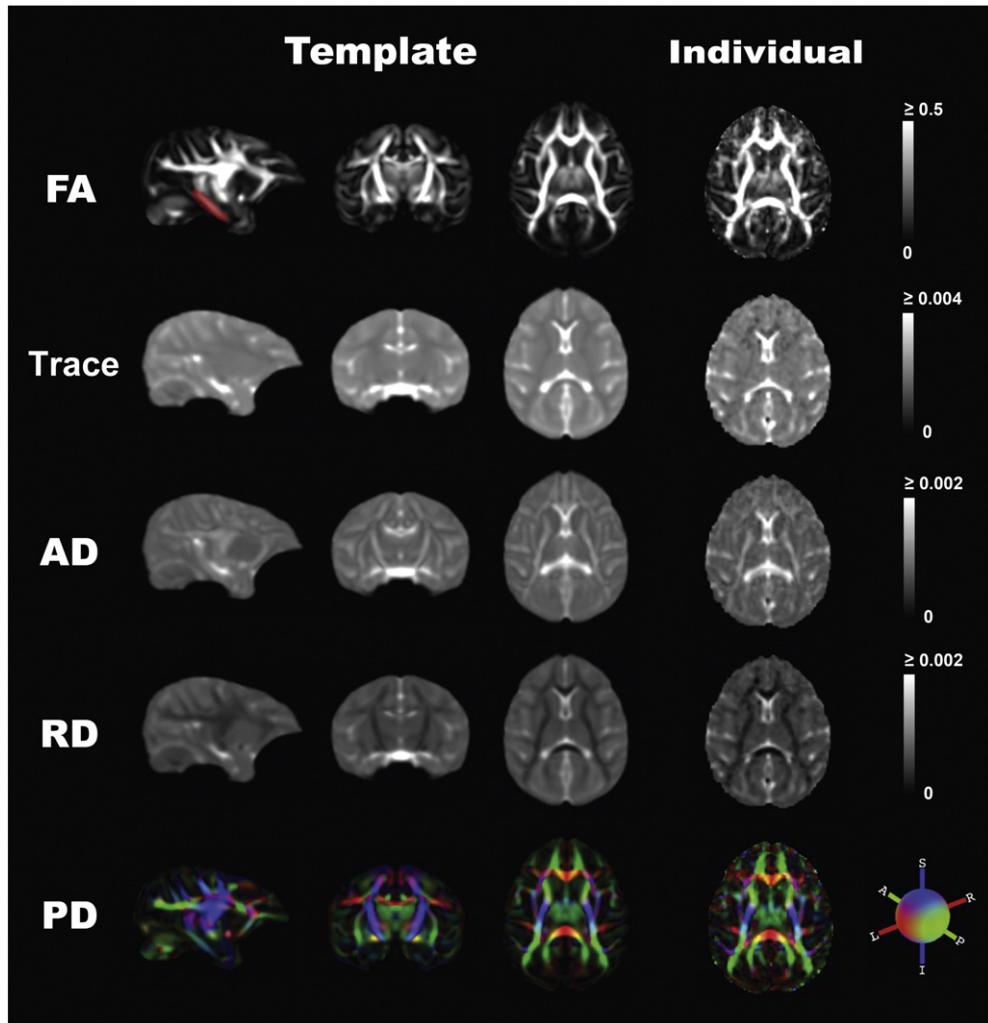


Fig. 3. DTI-based template. The first three columns show the properties of the final template, and the fourth column shows the properties of an individual monkey image. Fractional anisotropy (FA), trace, axial diffusivity (AD), radial diffusivity (RD) and principal diffusion direction images (PD) are presented in each row. The units of trace, AD, and RD are mm^2/s . The inferior cingulum mask required for tract-based spatial statistics (TBSS) is also shown on the sagittal slice of the FA image (in red color).

under the threshold of 0.2. An inferior cingulum mask was manually drawn on the template space to facilitate FA projection because the inferior cingulum tract had no well-defined search direction for the projection due to its tubular skeleton topology (Smith et al., 2006). After the projection, linear mixed-effect models were used to analyze the skeletonized FA images. Concurrently, TBSS of other diffusion-derived measures, including mean diffusivity (MD), axial diffusivity (AD) and radial diffusivity (RD) were also performed using the same approach.

Statistical analysis

For the total brain volume and ROI-based tissue volume, time-related change and PCP-related change were analyzed using linear mixed-effect models by the NLME package of the R language (Pinheiro et al., 2007). Main (fixed) effects of the time, the group factor (control, PCP1 and PCP2) and the interaction between the time and the group factors (time-group interaction) were included in the analysis; and the random effect (subject-specific intercept) of each monkey was also modeled to account for repeated measurements. Time could be modeled as a categorical factor to test the difference between different time points (M1), or it could be modeled as a continuous variable to test the linear/quadratic trend (M2), resulting in two similar but different models. For the M2, the quadratic term of time was not significant in any of the measures; thus, a linear fit was calculated. Besides, more complex structures of variance, correlation and random effect were also modeled, including heterogeneous variance, correlation structure with an autoregressive process of order 1, and random effect with both subject-specific intercept and slope. These models were compared by likelihood-ratio tests and information criteria (AIC and BIC), but none of these complex models outperformed the mixed-effect models M1 and M2. Thus, M1 and M2 were chosen for the final mixed-effect model analysis, which were followed by Tukey post-hoc tests for any significant factors.

Similar models (M1 and M2) were fitted for VBM and TBSS by SurfStat (Worsley, 2009), which was capable of fitting mass-univariate/multivariate linear mixed-effect models for the volumetric data. As the F tests for mixed effect models were not featured in SurfStat, different contrasts were designed to test different factors.

DBM was calculated from a nonlinear alignment between the images of two scans. For each pair of time points, a voxelwise one-sample permutation test was performed on the maps of Jacobian determinants to determine if there was a significant expansion or reduction of the brain using the *randomise* of FSL (Smith et al., 2004).

Results

Templates of infant monkeys

To avoid any bias to a particular time point and to improve the quality of templates, we created high-quality MRI and DTI templates using the data of all scans. The templates were a pre-requisite for the subsequent brain imaging analysis, and served as a target space to which individual brains were spatially normalized. Both the MRI-based template and the DTI-based template are provided in the Supplementary Data for downloading and usage.

MRI-based template

As illustrated in Fig. 2, the template had a superior contrast between different tissues and a better signal-to-noise ratio when compared to individual brains. The greater contrast and improved delineation of anatomical structures in the template demonstrated a good alignment of individual monkey brains. Moreover, the template contained brain parcellation maps derived from the UNC Primate Brain Atlas. Thus, we can use the template to label each brain region and to conduct ROI-based analysis.

DTI-based template

The properties of the DTI-based template and a representative individual example, including fractional anisotropy (FA), trace (trace = 3× mean diffusivity), axial diffusivity (AD), radial diffusivity (RD), and principal diffusion direction images (PD) are shown in Fig. 3. We also drew manually an inferior cingulum mask on the template space to facilitate its skeleton-projection for the following TBSS analysis. Like the MRI-based template, the DTI-based template had a better contrast than individual brains, which demonstrated a good alignment.

Brain development and the effect of PCP

We characterized the rapid development of the monkey brain and the influence of PCP during late infancy based on longitudinal brain imaging data. The results across S6, S10 and S16 are described in this section, as 1) the three time points showed evident changes in brain maturation, 2) S10 and S16 were far apart from the baseline scan (S6) and suffered less from the variability of the ages of different monkeys, 3) the three time points had all 14 monkeys scanned, which allowed more solid statistical inference.

Total brain volume

We calculated the total brain volume based on individual de-skinned structural images. Mixed-effect model analysis showed a significant growth of brain volume during late infancy ($F(2,22) = 154.63$, $p < 0.001$): on the average, the 10-month-old brain and the 16-month-old brain were $1.062 \times$ and $1.092 \times$ larger than the 6-month-old brain, respectively. However, the rate of brain growth gradually slowed down: an increase of 6.21% occurred from 6 to 10 months (adjusted $p < 0.001$), and was then followed by another increase of 2.81% from 10 to 16 months (adjusted $p < 0.001$) (Fig. 4).

Although the PCP-treated groups had higher growth rates (11.15% in PCP1, 8.86% in PCP2) than the control group (7.13%) from 6 to 16 months (Fig. 4), neither the group factor ($F(2,11) = 0.38$, $p = 0.69$) nor the time-group interaction ($F(4,22) = 2.58$, $p = 0.07$) had significant effects on the changes of total brain volume.

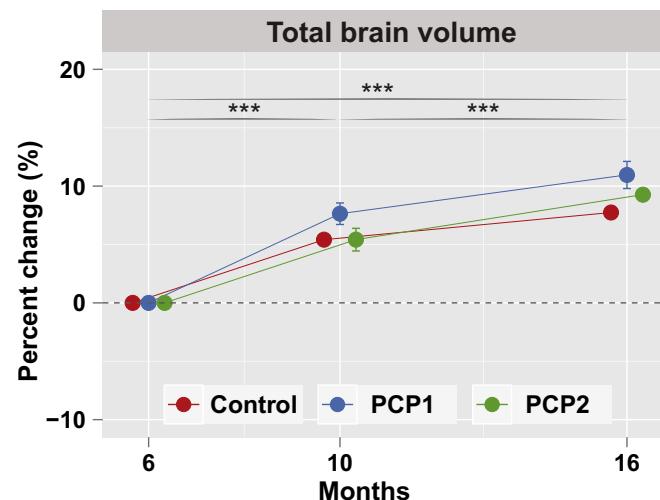


Fig. 4. Total brain volume changes. The percent changes of total brain volume of each scan against the baseline scan (S6) are presented in mean \pm SEM for each group. Mixed-effect model analysis (based on absolute volume) shows significant time-related changes ($F(2,22) = 154.63$, $p < 0.001$), but non-significant effects of the group factor ($F(2,11) = 0.38$, $p = 0.69$) and non-significant time-group interaction ($F(4,22) = 2.58$, $p = 0.07$). Asterisks indicate the difference of volume between the scans based on post-hoc tests: * $p < 0.05$, ** $p < 0.01$, and *** $p < 0.001$. Detailed statistics of mixed-model analyses are presented in Table S2.

ROI-based tissue volume analysis

Development of the monkey brain during late infancy was characterized by reduced gray matter volume and increased white matter volume (Figs. 5 and 6). However, the developmental trend of tissue volume was regionally specific. Gray matter volume in the occipital lobe, cerebellum, and insular cortex decreased significantly after S6. Gray matter in other cortices, including the frontal, parietal, and temporal lobes, and cingulate cortex, decreased significantly only after S10. White matter volume displayed an overall increasing trend in most brain areas, but not in the occipital lobe ($F(2,22) = 1, p = 0.38$). A non-significant increase of white matter volume was also found in the parietal lobe ($F(2,22) = 2.52, p = 0.10$). The increase of white matter volume from S6 to S10 was significant in the cerebellum and corpus callosum, but not significant in the frontal lobe, temporal lobe, and cingulate cortex.

Although neither the group factor nor the time-group interaction had significant effects on any of the above changes, the PCP groups displayed a non-significant acceleration of the decrease in gray matter volume and of the increase in white matter volume after S10 in most brain regions, including the frontal, parietal, and temporal lobes, cerebellum, and cingulate cortex (Figs. 5 and 6). All detailed statistics of mixed-model analyses and the absolute volume of the above measures are presented in Table S2.

Voxel-based morphometry (VBM) and Deformation-based morphometry (DBM)

ROI-based analysis provided measures of large areas, whereas VBM enabled us to compare changes of tissue volume at the voxel level. Consistent with ROI-based analysis, VBM revealed reduced gray matter

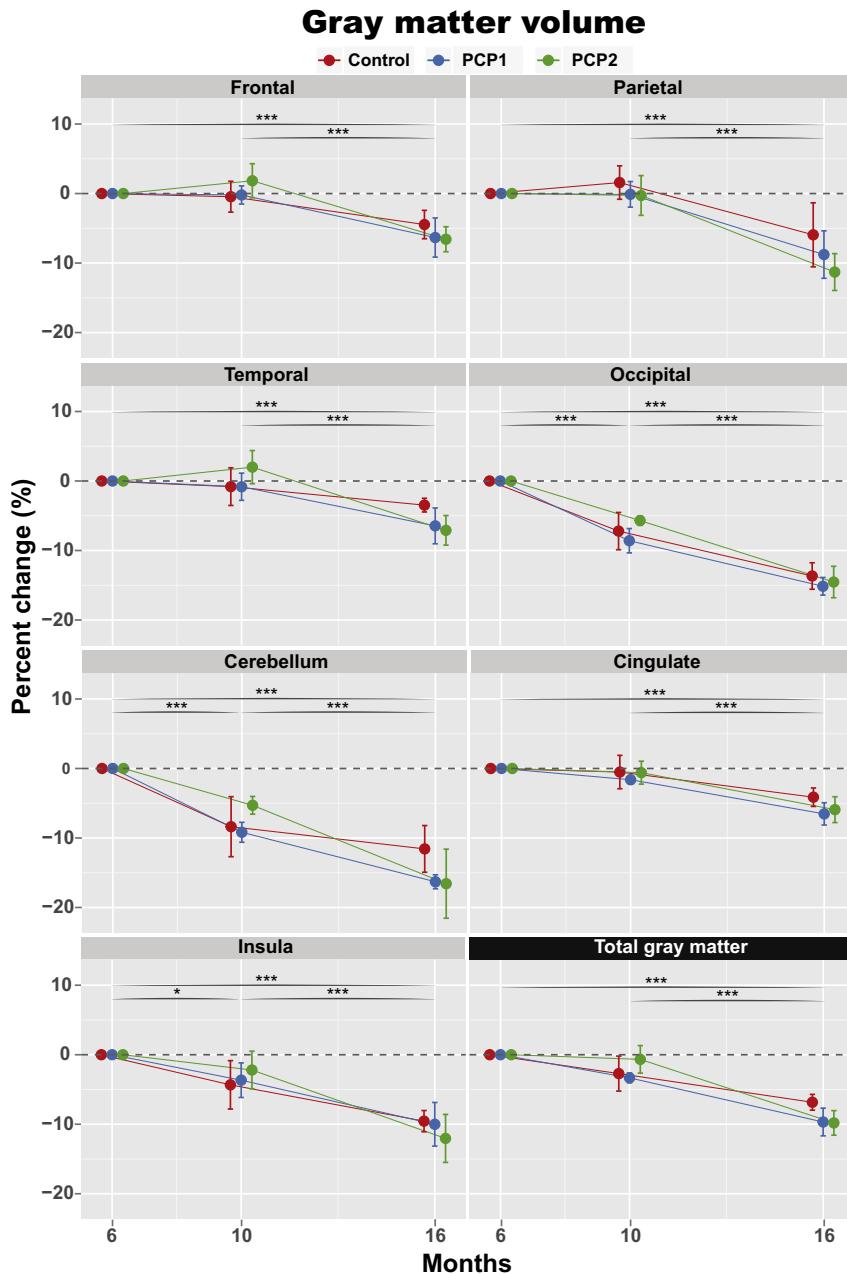


Fig. 5. ROI-based gray matter volume analysis. Percent changes of gray matter volume of each scan against the baseline scan (S6) are presented in mean \pm SEM for each group. Mixed-effect model analysis (based on absolute volume) shows significant time-related changes in all brain regions. Neither the group factor nor the time-group interaction is significant in any brain regions. Asterisks indicate differences of volume between the scans based on post-hoc tests: * $p < 0.05$, ** $p < 0.01$, and *** $p < 0.001$. Detailed statistics of mixed-model analyses are presented in Table S2.

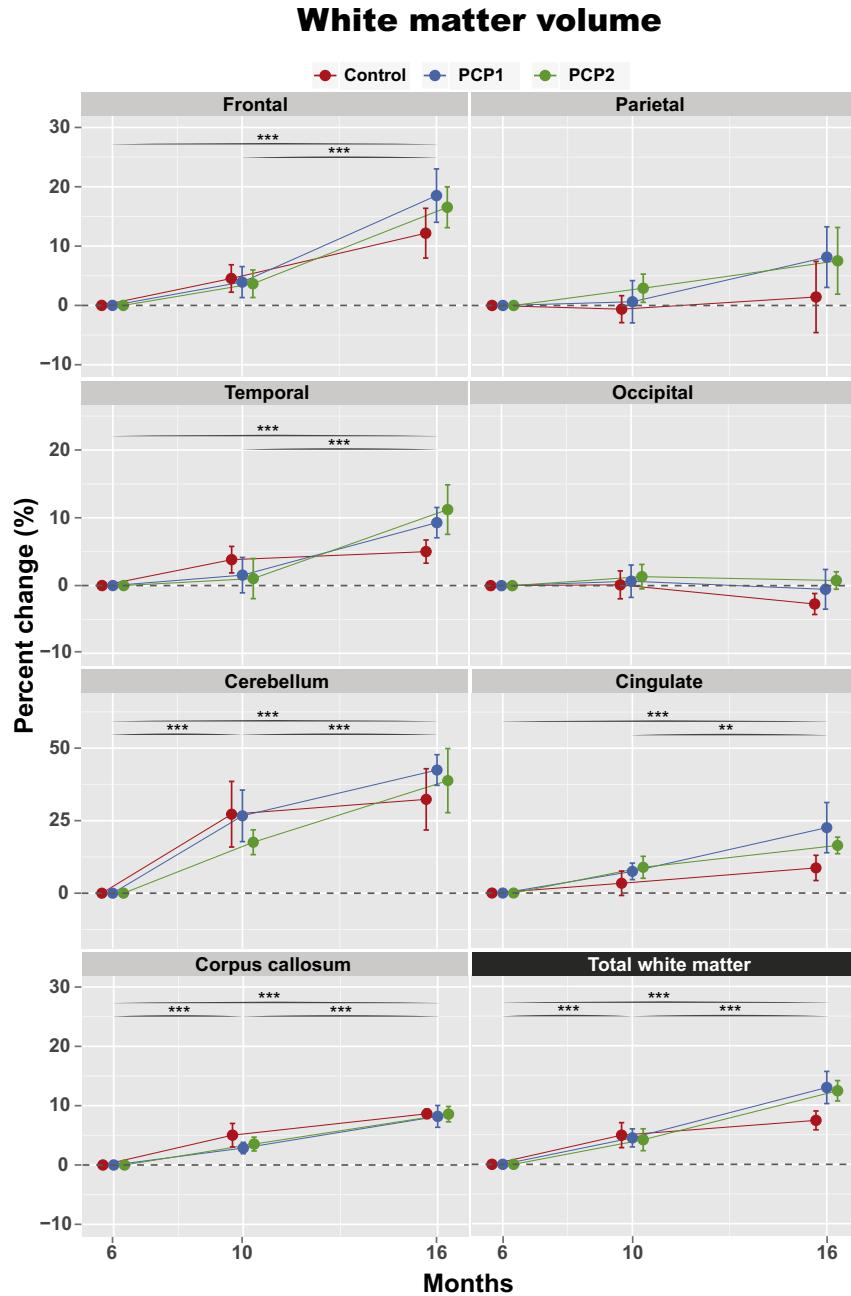


Fig. 6. ROI-based white matter volume analysis. Percent changes of white matter volume of each scan against the baseline scan (S6) are presented in mean \pm SEM for each group. Mixed-effect model analysis (based on absolute volume) shows significant time-related changes in most brain regions, but not in the occipital and parietal lobes. Neither the group factor nor the time-group interaction is significant in any brain region. Asterisks indicate the difference of volume between the scans based on post-hoc tests: * $p < 0.05$, ** $p < 0.01$, and *** $p < 0.001$. Detailed statistics of mixed-model analyses are presented in Table S2.

volume and increased white matter volume across the brain during late infancy (Fig. S1). VBM also showed regional variations: white matter changes in the occipital and parietal lobes were less obvious than the changes in other brain regions. Neither the group factor nor the time-group interaction had significant effect in the changes of VBM.

DBM was an effective technique to detect morphological differences based on the deformation field derived from nonlinear alignments. Consistent with the above findings, more expansion was found in the white matter than the gray matter, whereas more reduction was found in the gray matter than the white matter (Fig. 7).

Tract-Based Spatial Statistics (TBSS)

Besides volume and shape changes, we characterized changes in white matter microstructure based on DTI. TBSS showed increased

fractional anisotropy (FA) and decreased diffusivity (MD, RD and AD) across all major tracts of the brain during late infancy (Fig. 8). The decrease in RD (diffusivity perpendicular to the axon) was more evident than that of AD (diffusivity along the axon). Thus, changes in FA and MD were driven mainly by the reduction in RD. All of these changes were more obvious at S16 than S10, suggesting continuous development of the white matter microstructure during late infancy. None of the measures showed any significant effect of PCP on brain development.

Discussion

This study demonstrates that 1) the volume of gray matter decreases in monkeys during late infancy, much earlier than in humans; 2) the volume of white matter increases and is associated with an increase in

Deformation-based morphometry

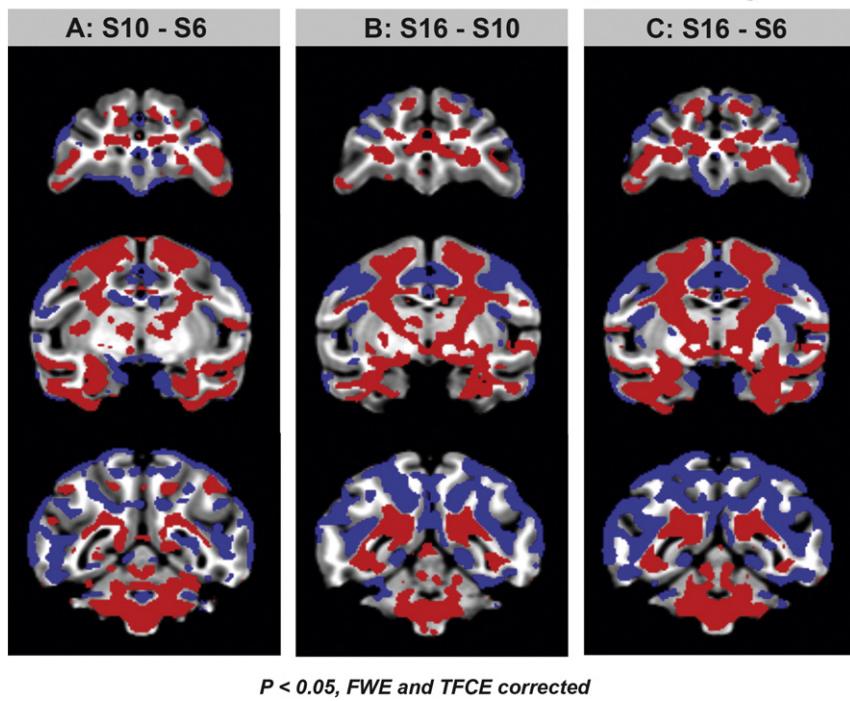


Fig. 7. Deformation-based morphometry. The red color represents significant expansion and the blue color represents significant reduction based on the permutation test implemented in the *randomise* of FSL (Smith et al., 2004).

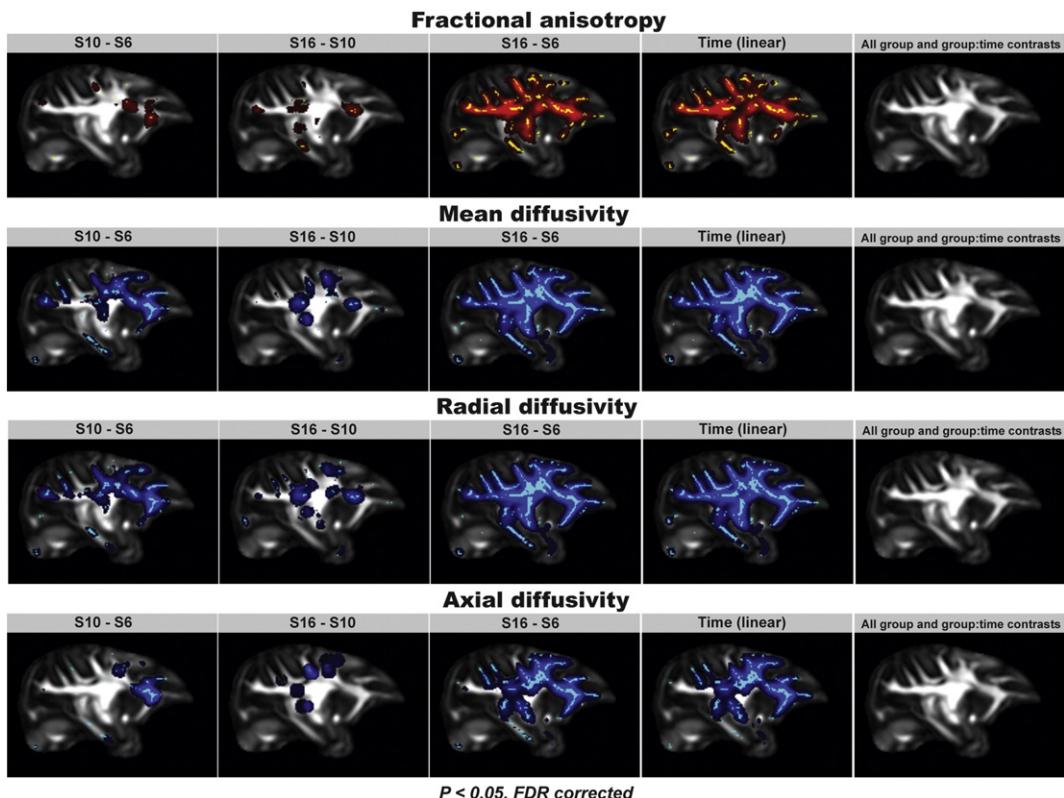


Fig. 8. Tract-Based Spatial Statistics (TBSS). Fractional anisotropy, mean diffusivity, axial diffusivity, and radial diffusivity are analyzed by mass-univariate linear mixed-effect models. The first three figures of each property represent the time-related contrasts of M1 where the time is modeled as categorical factors; the fourth figure represents the time-related contrasts of M2 where the time is modeled as a continuous variable; and the fifth figure represents all contrasts related to the group factor and the time-group interaction. All skeletonized results (yellow and azure) are filled into local tracts (red and blue) by the *tbss_fill* of FSL for a better visualization (Smith et al., 2006). The red and yellow colors represent significant increases, and the blue and azure colors represent significant decreases in these properties.

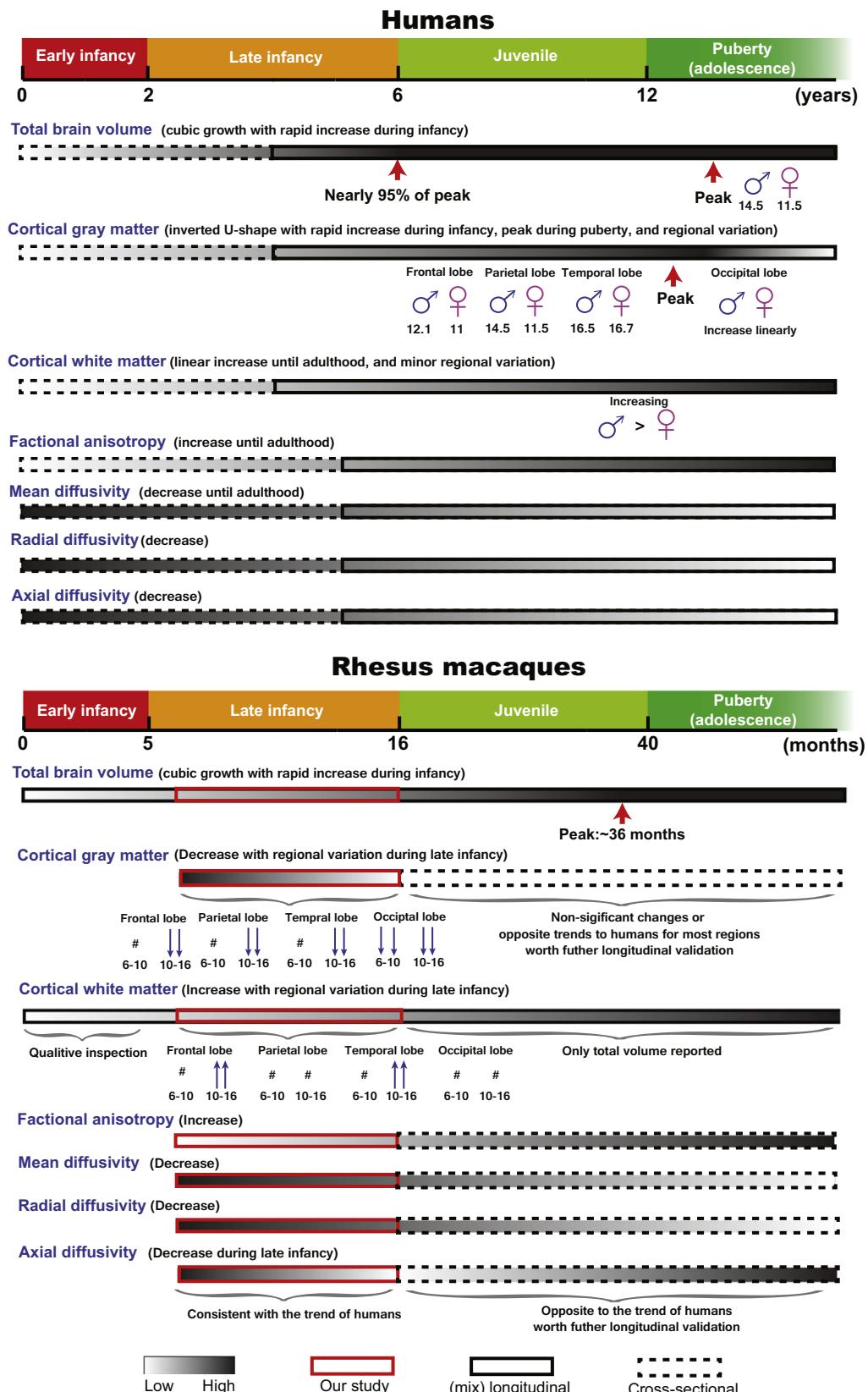


Fig. 9. Summary of human and monkey brain development from early infancy to puberty (adolescence). Adopted from a recent study (Sakai et al., 2011), these developmental stages are defined based on combined dental eruption and sex maturation for inter-specific comparisons (Plant, 1994, 2001; Plant and Barker Gibb, 2004; Smith et al., 1994; Terasawa and Fernandez, 2001). The time scale of monkeys is re-scaled to match that of humans for a better comparison. These developmental trajectories are depicted mainly based on longitudinal and mixed-longitudinal data (Brouwer et al., 2012; Giedd, 2004; Giedd et al., 1999; Lebel and Beaulieu, 2011; Malkova et al., 2006) with the cross-sectional data as supplements (Geng et al., 2012; Knickmeyer et al., 2008, 2010; Lenroot et al., 2007; Shi et al., 2013). The phases based on (mixed-)longitudinal data are marked with solid outlines, the cross-sectional data with dashed outlines, and our longitudinal data with red solid outlines. The color bar from white to black represents an increase in measures, and vice versa.

fractional anisotropy and a decrease in diffusivity; and 3) the maturation of gray and white matter are regionally specific; and 4) PCP tends to accelerate maturation after 10 months of age, although not statistically significant.

We observed a rapid growth of monkey brain volume during early development, consistent with a previous study (Malkova et al., 2006). However, the growth rate began to slow down during late infancy, as 6.21% of the expansion occurred between 6 and 10 months and only 2.81% between 10 and 16 months (Fig. 4). The fast expansion is mainly caused by the significant increase of white matter volume (Fig. 6), while the later decline in the expansion rate can be partly explained by a significant decrease of gray matter volume after 10 months (Fig. 5). Compared with monkey brain development, human brain volume increases 100% during the first year of life, and then keeps cubic growth into adolescence (Fig. 9) (Giedd, 2004; Giedd et al., 1999; Gilmore et al., 2007), which indicates that humans have a less mature brain at birth and a longer period of early rapid brain growth than monkeys (Gibson and Petersen, 1991).

The changes in the volume of gray and white matter volume of monkey brains were regionally specific (Figs. 5 and 6). The occipital lobe and cerebellum (presumably important for primary functions) had a significant decrease in gray matter volume before 10 months, whereas the frontal, temporal, and parietal lobes (associated with functional integrations) had significant decreases only after 10 months. As to white matter, its volume showed an increasing trend in most brain regions during late infancy, but not in the occipital lobe. The developmental pattern demonstrates that brain regions for primary functions mature earlier than brain regions associated with functional integration. In humans, brain maturation shows more complex regional variations (Fig. 9). Gray matter volume in brain regions for functional integration, including the frontal, temporal, and parietal lobes, decreases only after adolescence and the change tends to follow a trajectory of an “inverted U curve” with peaks at different times, whereas the gray matter of the occipital lobe keeps increasing without significant decline or leveling (Giedd, 2004; Giedd et al., 1999). Compared with monkeys, the more complex and delayed brain development of humans allows a stronger influence of postnatal experiences on brain structures, which may contribute to the emergence of human-specific cognitive abilities and social behaviors.

In addition to volume change, we also characterized the development of white matter microstructure using DTI. We observed an overall increase in FA and a decrease in diffusivity (MD, RD, and AD). Previous studies revealed increased RD and decreased FA in mice that were genetically modified or treated with cuprizone to be myelin-deficient (Nair et al., 2005; Song et al., 2002, 2005). As demyelination contributes to a decrease of FA and an increase of RD, we infer that myelination may be the main reason for the change of FA and RD during late infancy. We also observed a decrease in AD, which is consistent with previous findings in humans (Fig. 9) (Forbes et al., 2002; Mukherjee et al., 2001; Sadeghi et al., 2013; Schneider et al., 2004; Zhang et al., 2005). Contrary to human development, a previous cross-sectional DTI study in monkeys showed an increase of AD from late infancy to young adulthood (Shi et al., 2013). The above evidence indicates that the change of AD in monkeys may be nonlinear after late infancy, which is worth further longitudinal validation (Fig. 9).

We also investigated the effect of PCP on these trajectories with the hope of using monkeys as an animal model for schizophrenia (Jones et al., 2011; Mouri et al., 2007). Despite their trajectories being similar to those of the control group, the PCP-treated groups had an accelerated rate of development after 10 months (Figs. 4–6). However, none of the PCP-related changes was significant. Two possible reasons may account for the non-significance. The first may be the small sample size (14 monkeys). However, a longitudinal study like ours has a higher power and requires fewer samples than cross-sectional and mixed-longitudinal studies. In addition, we are mainly interested in major PCP-treatment effects that are detectable even with a small sample

size. The second reason may be that the time of PCP treatment was beyond the critical phases of the development of the glutamatergic circuitry in the prefrontal cortex that PCP primarily affects: one is from late gestation to the first 3 months postnatal, when the number of excitatory synapses (glutamatergic) substantially increases; the other from around 15–18 months to young adulthood, when the number of excitatory synapses begins to decline until the stable adult level is achieved (Hoftman and Lewis, 2011). In our study, the time points of PCP treatment and scans were within the stable period from 3 to 15–18 months, which partly explains the non-significant effect of PCP. Therefore, it is worthwhile to study if the effect of PCP becomes significant or disappears when these monkeys reach adulthood, and whether these changes can influence cognitive performance that are demanding for infant monkeys.

Besides the issue of PCP, we cannot investigate sex differences in brain development because the monkeys involved in our study were all males. Although the general developmental trajectory of males may be applicable to females, gender differences in early brain development may exist and are worth future investigation. Furthermore, our longitudinal data only cover development during late infancy, a time scale that needs to be expanded in the future.

Supplementary data to this article can be found online at <http://dx.doi.org/10.1016/j.neuroimage.2014.11.056>.

Acknowledgments

This work was supported by 973 program (2012CB825500, 2011CB707800), NSFC (31271168, 91132307/H09, 61178051) and NSFC Finnish-Chinese joint project 813111172. The authors would like to thank Prof. Ziad Hafed, Prof. Synnöve Carlson, Dr. Ilkka Linnankoski and Joshua Rizak for reading and revising the manuscript.

Conflict of interest

None declared.

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